

Bower, L., Jones, T., Ban, J., Chen, H., Cheuk, R., Chung, M. K.,
 Kim, C., Lin, J., Liu, S. X., Pham, P. K., Sakano, H., Shin, P.,
 Yamada, K., Ecker, J., Theologis, A. and Davis, R. W.
 Nguyen, M. (USP/Stanford) and Seki, M. (RIKEN GSC) contributed
 equally to this work. Shirokai, K. (RIKEN GSC) and Davis, R. W.
 (USP/Stanford) contributed equally to this work as PIs.
 Location/Qualifiers
 1. .566
 /organism="Arabidopsis thaliana"
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 /chromosome="5"
 /note="AF109-78-L12"
 /note="This clone is in plasmid script
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 137 c 149 g 199 t

Query Match 5.9% Score 173; DB 1; Length 666;
 Best Local Similarity 72.7%; Pred. No. 0.73;
 Matches 209; Conservative 0; Mismatches 60; Indels 0; Gaps 0;

QY 58 AAGCGCTTGAAGTGAAGAAAGTGAATGACATGACCTGCTGGCTGGATATGCTT 117
 DB 175 AAGCGCTTGAAGTGAAGAAAGTGAATGACATGACCTGCTGGCTGGATATGCTT 234
 QY 118 GATTAAGTGCATTCGATCGACGACACCAATTAATGATCTTTTCATGAAATGCAAGTAAAC 177
 DB 235 GATTAAGTGCATTCGATCGACGACACCAATTAATGATCTTTTCATGAAATGCAAGTAAAC 294
 QY 178 CAGGCGTCCGCTACTTCAGAGAGTGTACTGTGCATGGAGATGTGTACCAATGCTTTT 237
 DB 295 CAGGCGTCCGCTACTTCAGAGAGTGTACTGTGCATGGAGATGTGTACCAATGCTTTT 354
 QY 238 CACTTCACATGATCTCTGCTGCTGCTCAAAAGCAGCAGGTGTGCATTTGGACACAGA 297
 DB 355 CACTTCACATGATCTCTGCTGCTGCTCAAAAGCAGCAGGTGTGCATTTGGACACAGA 414
 QY 298 GACTGGGAATTCGAAAAGTATGCTGACACTA 326
 DB 415 GACTGGGAATTCGAAAAGTATGCTGACACTA 443

RESULT 3
 SCYOL134C 1255 bp DNA linear PLN 11-AUG-1997
 LOCUS S. cerevisiae chromosome XV reading frame ORF YOL134C.
 DEFINITION 274876 Y13140
 ACCESSION 274876.1 GI:1420022
 KEYWORDS
 SOURCE Saccharomyces cerevisiae.
 ORGANISM Saccharomyces cerevisiae.
 Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
 1 (bases 1 to 1255)
 AUTHORS Arino, J., Casamayor, A., Cano, F. J., Gancedo, C., Lafuente, M. J.,
 Aldea, M., Cases, C. and Herrero, E.
 JOURNAL Published
 REFERENCE 2 (bases 1 to 1255)
 TITLE Direct Submission
 AUTHORS Submitted (04-JUL-1996) data collected by MIPS on behalf of the
 JOURNAL European yeast chromosome XV sequencing project. MIPS at the

FEATURES
 source
 Max-Planck-Institut fuer Biochemie, Am Klopferspitz 18a D-82152
 Martinsried, FRG; E-mail: Mews@mpi-bpc.mpg.de
 Location/Qualifiers
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 137 c 149 g 199 t

BASE COUNT 419 a 209 c 249 g 378 t

Query Match 37.8% Score 123.6; DB 3; Length 1255;
 Best Local Similarity 66.5%; Pred. No. 0.84;
 Matches 177; Conservative 0; Mismatches 89; Indels 0; Gaps 0;

QY 55 AAGAGCGCTTGAAGTGAAGAAAGTGAATGACATGACCTGCTGGCTGGATATGCTT 114
 DB 374 AAGAGCGCTTGAAGTGAAGAAAGTGAATGACATGACCTGCTGGCTGGATATGCTT 433
 QY 115 GATTAAGTGCATTCGATCGACGACACCAATTAATGATCTTTTCATGAAATGCAAGTAAAC 174
 DB 434 GATTAAGTGCATTCGATCGACGACACCAATTAATGATCTTTTCATGAAATGCAAGTAAAC 493
 QY 175 CAGGCGTCCGCTACTTCAGAGAGTGTACTGTGCATGGAGATGTGTACCAATGCTTTT 234
 DB 494 CAGGCGTCCGCTACTTCAGAGAGTGTACTGTGCATGGAGATGTGTACCAATGCTTTT 553
 QY 235 TTTCACTTCGATGATCTCTGCTGCTGCTCAAAAGCAGCAGGTGTGCATTTGGACACAGA 294
 DB 554 TTTCACTTCGATGATCTCTGCTGCTGCTCAAAAGCAGCAGGTGTGCATTTGGACACAGA 613
 QY 295 AGACGATGCGAATTCGAAAAGTATGCTGACACTA 320
 DB 614 AGACGATGCGAATTCGAAAAGTATGCTGACACTA 639

RESULT 4
 AY052401 357 bp mRNA linear PLN 16-OCT-2001
 LOCUS Arabidopsis thaliana ring box-1-like protein mRNA, complete cds.
 DEFINITION AY052401
 ACCESSION AY052401.1 GI:16186264
 KEYWORDS
 SOURCE Arabidopsis thaliana.
 ORGANISM Arabidopsis thaliana.
 Eukaryota; Eudicotyledons; Streptophyta; Embryophyta; Tracheophyta;
 Spermatophyta; Angiosperms; Eudicotyledons; Core eudicots;
 Rosidae; eudicots II; Brassicales; Brassicaceae; Arabidopsis.
 1 (bases 1 to 357)
 AUTHORS O'Keefe, L.
 JOURNAL Direct Submission
 REFERENCE 25-AUG-2001 Institute of Plant Biology, Biological
 Research Center, Temesvári krt. 62, Szeged H-6701, Hungary
 TITLE Submitted (25-AUG-2001) data collected by MIPS on behalf of the
 JOURNAL European yeast chromosome XV sequencing project. MIPS at the

FEATURES
 source
 Max-Planck-Institut fuer Biochemie, Am Klopferspitz 18a D-82152
 Martinsried, FRG; E-mail: Mews@mpi-bpc.mpg.de
 Location/Qualifiers
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 137 c 149 g 199 t

CDS

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BASE COUNT 91 a 83 c 94 g 89 t

Query Match 4.0% Score 13; DB 2; Length 357;
Best Local Similarity 65.5% Pred. No. 11;
Matches 19; Conservative 0; Mismatches 10; Indels 0; Gaps 0;

OR 22 GATATCCCAAGCCGACACAGACGCGCC 50
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Db 141 GATATCCCAAGCCGACAGACGCGCC 113

RESULT 5
LOCUS AY072430/c 666 bp mRNA linear PLN 21-JAN-2002
DEFINITION Arabidopsis thaliana ring-box protein - like (At5g20570) mRNA.
ACCESSION AY072430
VERSION AY072430.1 GI:18252990
KEYWORDS FIL CDNA.
SOURCE Arabidopsis thaliana.
ORGANISM Arabidopsis thaliana.
REFERENCE 1. Kawai, Y., Kim, C., Lin, J., Liu, S.X., Natunaka, M., Ishida, J., Kawai, A., Sakurai, T., Satou, M., Seki, M., Shinozaki, K., Shinozaki, K., Ecker, J., Theologis, A. and Davis, R.W. Direct Submission Submitted (02-JAN-2002) DNA Sequencing and Technology Center, Stanford University, 855 California Avenue, Palo Alto, CA 94304, USA
COMMENT e-mail for correspondence: arabis@sequence.stanford.edu

RIKEN Genomic Sciences Center (GSC) members carried out the collection and clustering of RFLP CDNA (RAFL CDNA : 'RIKEN Arabidopsis Full-length CDNA') : Seki, M., Natunaka, M., Ishida, J., Satou, M., Kamiya, A., Sakurai, T., Carninci, P., Kawai, J., Hayashizaki, Y. and Shinozaki, K.

The Salk, Stanford, RGC (SSP) Consortium members carried out the sequencing and annotation of the RFLP CDNA : Nguyen, M., Soultzer, L., Jones, T., Banb, J., Chen, H., Cheuk, R., Kim, C., Lin, J., Liu, S.X., Pham, P.K., Sakano, H., Shinozaki, K., Ecker, J., Theologis, A. and Davis, R.W.

Nguyen, M. (SSP/Stanford) and Seki, M. (RIKEN GSC) contributed equally to this work. Shinozaki, K. (RIKEN GSC) and Davis, R.W. (SSP/Stanford) contributed equally to this work as PIs.

FEATURES

SOURCE

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CDS

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BASE COUNT 181 a 137 c 149 g 199 t

Query Match 4.0% Score 13; DB 1; Length 666;
Best Local Similarity 65.5% Pred. No. 5; 7;
Matches 19; Conservative 0; Mismatches 10; Indels 0; Gaps 0;

OR 22 GATATCCCAAGCCGACACAGACGCGCC 50
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Db 228 GATATCCCAAGCCGACAGACGCGCC 200

RESULT 5
LOCUS SCYOL134C 1255 bp DNA linear PLN 11-NOV-1997
DEFINITION S.cerevisiae chromosome XV reading frame ORF YOL134C.
ACCESSION Z74876.1
VERSION Z74876.1 GI:1420022
KEYWORDS
SOURCE Saccharomyces cerevisiae.
ORGANISM Saccharomyces cerevisiae.
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